

Single-cell RNA sequencing of endosteal bone marrow cells

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Single cell transcriptomics identifies a unique adipose lineage cell population that regulates bone marrow environment

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1. Zhong, L. (2020). Single-cell RNA sequencing of endosteal bone marrow cells. Bio-protocol Preprint. bio-protocol.org/2020/06/01/prep601.
2. Zhong, L., Yao, L., Tower, R. J., Wei, Y., Miao, Z., Park, J., Shrestha, R., Wang, L., Yu, W., Holdreith, N., Huang, X., Zhang, Y., Tong, W., Gong, Y., Ahn, J., Susztak, K., Dymment, N., Li, M., Long, F., Chen, C., Seale, P. and Qin, L.(2020). Single cell transcriptomics identifies a unique adipose lineage cell population that regulates bone marrow environment. eLIFE. DOI: [10.7554/eLife.54695](https://doi.org/10.7554/eLife.54695)

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